

PA0394 PCT - Sequence Listing - PatentIn 3.2.ST25  
SEQUENCE LISTING

<110> Amersham Biosciences UK Limited

<120> Cytochrome C Protein and Assay

<130> PA0394

<140> GB0329353.7

<141> 2003-12-19

<160> 15

<170> PatentIn version 3.2

<210> 1

<211> 315

<212> DNA

<213> Homo sapiens

<400> 1

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acaggtcagg cccctggata ctcttacaca gccgccaata agaacaaagg catcatctgg      180
ggagaggata cactgatgga gtatttggag aatcccaaga agtacatccc tggaacaaaa      240
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aaagctacta atgag                                     315

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<210> 2

<211> 105

<212> PRT

<213> Homo sapiens

<400> 2

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Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr Gly Pro Asn
20          25          30

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Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro Gly Tyr Ser
35          40          45

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Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly Glu Asp Thr
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Leu Met Glu Tyr Leu Glu Asn Pro Lys Lys Tyr Ile Pro Gly Thr Lys
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Met Ile Phe Val Gly Ile Lys Lys Lys Glu Glu Arg Ala Asp Leu Ile
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## PA0394 PCT - Sequence Listing - Patentin 3.2.ST25

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 aaacttacc ttaaatttat ttgcactact ggaaaactac ctgttccatg gccaacactt 180  
 gtcactactc tctcttatgg tgttcaatgc ttttcaagat acccagatca tatgaaacgg 240  
 catgactttt tcaagagtgc catgcccgaagggttatgtac aggaaagaac tatatttttc 300  
 aaagatgacg ggaactacaa gacacgtgct gaagtcaagt ttgaagggtga tacccttggt 360  
 aatagaatcg agttaaaagg tattgatattt aaagaagatg gaaacattct tggacacaaa 420  
 ttggaataca actataactc acacaatgta tacatcatgg cagacaaaca aaagaatgga 480  
 atcaaagtta acttcaaaat tagacacaac attgaagatg gaggcgttca actagcagac 540  
 cattatcaac aaaatactcc aattggcgat ggccctgtcc ttttaccaga caaccattac 600  
 ctgtccacac aatctgccct ttcgaaagat cccaacgaaa agagagacca catggtcctt 660  
 cttggctttg taacagctgc tgggattaca catggcatgg atgaactata caaactcgag 720  
 aattcgacca tgggtgatgt tgagaaaggc aagaagattt ttattatgaa gtgttcccag 780  
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<220>  
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 Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu  
 20 25 30  
 Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys  
 35 40 45

## PA0394 PCT - Sequence Listing - Patentin 3.2.ST25

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu  
 50 55 60

Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg  
 65 70 75 80

His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg  
 85 90 95

Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val  
 100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile  
 115 120 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn  
 130 135 140

Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly  
 145 150 155 160

Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val  
 165 170 175

Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro  
 180 185 190

Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser  
 195 200 205

Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Gly Phe Val  
 210 215 220

Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Leu Glu  
 225 230 235 240

Asn Ser Thr Met Gly Asp Val Glu Lys Gly Lys Lys Ile Phe Ile Met  
 245 250 255

Lys Cys Ser Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr  
 260 265 270

Gly Pro Asn Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro  
 275 280 285

Gly Tyr Ser Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly  
 290 295 300

Glu Asp Thr Leu Met Glu Tyr Leu Glu Asn Pro Ala Lys Tyr Ile Pro  
 305 310 315 320

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Gly Thr Lys Met Ile Phe Val Gly Ile Lys Lys Lys Glu Glu Arg Ala  
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<220>  
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 acaggtcagg cccctggata ctcttacaca gccgccaata agaacaaagg catcatctgg 180  
 ggagaggata cactgatgga gtatttggag aatcccgcca agtacatccc tggaacaaaa 240  
 atgatctttg tcggcattaa gaagaaggaa gaaagggcag acttaatagc ttatctcaaa 300  
 aaagctacta atgaggggtcg acccgggatg agtaaaggag aagaactttt cactggagtt 360  
 gtccaatttc ttgttgaatt agatggtgat gttaatgggc acaaattttc tgtcagtgga 420  
 gaggggtgaag gtgatgcaac atacggaaaa cttaccctta aatttatttg cactactgga 480  
 aaactacctg ttccatggcc aacacttgtc actactctct cttatggtgt tcaatgcttt 540  
 tcaagatacc cagatcatat gaaacggcat gactttttca agagtgccat gcccgaaggt 600  
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 <213> Artificial

<220>  
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<400> 6

Met Gly Asp Val Glu Lys Gly Lys Lys Ile Phe Ile Met Lys Cys Ser  
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## PA0394 PCT - Sequence Listing - Patent in 3.2.ST25

Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr Gly Pro Asn  
 20 25 30  
 Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro Gly Tyr Ser  
 35 40 45  
 Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly Glu Asp Thr  
 50 55 60  
 Leu Met Glu Tyr Leu Glu Asn Pro Ala Lys Tyr Ile Pro Gly Thr Lys  
 65 70 75 80  
 Met Ile Phe Val Gly Ile Lys Lys Lys Glu Glu Arg Ala Asp Leu Ile  
 85 90 95  
 Ala Tyr Leu Lys Lys Ala Thr Asn Glu Gly Arg Pro Gly Met Ser Lys  
 100 105 110  
 Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp  
 115 120 125  
 Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly  
 130 135 140  
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 145 150 155 160  
 Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu Ser Tyr Gly  
 165 170 175  
 Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg His Asp Phe  
 180 185 190  
 Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe  
 195 200 205  
 Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu  
 210 215 220  
 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys  
 225 230 235 240  
 Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser  
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 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val  
 260 265 270  
 Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala  
 275 280 285

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Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu  
 290 295 300

Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser Lys Asp Pro  
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Asn Glu Lys Arg Asp His Met Val Leu Leu Gly Phe Val Thr Ala Ala  
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Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys  
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 gtcactactc tctcttatgg tggttcaatgc ttttcaagat acccagatca tatgaaacgg 240  
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 aaagatgacg ggaactacaa gacacgtgct gaagtcaagt ttgaagggtga tacccttggt 360  
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<210> 8  
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 <213> Artificial

## PA0394 PCT - Sequence Listing - Patentin 3.2.ST25

&lt;220&gt;

&lt;223&gt; synthetic polypeptide

&lt;400&gt; 8

Met Ser Lys Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val  
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Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu  
20 25 30

Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys  
35 40 45

Thr Thr Gly Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu  
50 55 60

Ser Tyr Gly Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg  
65 70 75 80

His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg  
85 90 95

Thr Ile Phe Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val  
100 105 110

Lys Phe Glu Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile  
115 120 125

Asp Phe Lys Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn  
130 135 140

Tyr Asn Ser His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly  
145 150 155 160

Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val  
165 170 175

Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro  
180 185 190

Val Leu Leu Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser  
195 200 205

Lys Asp Pro Asn Glu Lys Arg Asp His Met Val Leu Leu Gly Phe Val  
210 215 220

Thr Ala Ala Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys Leu Glu  
225 230 235 240

Asn Ser Thr Met Gly Asp Val Glu Lys Gly Lys Lys Ile Phe Ile Met  
245 250 255



## PA0394 PCT - Sequence Listing - Patent in 3.2.ST25

Lys Cys Ser Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr  
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 Gly Pro Asn Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro  
                   275                  280                  285  
 Gly Tyr Ser Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly  
                   290                  295                  300  
 Glu Asp Thr Leu Met Glu Tyr Leu Glu Asn Pro Lys Lys Tyr Ile Pro  
                   305                  310                  315                  320  
 Gly Thr Lys Met Ile Phe Val Gly Ile Lys Lys Lys Glu Glu Arg Ala  
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 Asp Leu Ile Ala Tyr Leu Lys Lys Ala Thr Asn Glu  
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PA0394 PCT - Sequence Listing - Patentin 3.2.ST25  
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1 5 10 15

Gln Cys His Thr Val Glu Lys Gly Gly Lys His Lys Thr Gly Pro Asn  
20 25 30

Leu His Gly Leu Phe Gly Arg Lys Thr Gly Gln Ala Pro Gly Tyr Ser  
35 40 45

Tyr Thr Ala Ala Asn Lys Asn Lys Gly Ile Ile Trp Gly Glu Asp Thr  
50 55 60

Leu Met Glu Tyr Leu Glu Asn Pro Lys Lys Tyr Ile Pro Gly Thr Lys  
65 70 75 80

Met Ile Phe Val Gly Ile Lys Lys Lys Glu Glu Arg Ala Asp Leu Ile  
85 90 95

Ala Tyr Leu Lys Lys Ala Thr Asn Glu Gly Arg Pro Gly Met Ser Lys  
100 105 110

Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp  
115 120 125

Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly  
130 135 140

Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly  
145 150 155 160

Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Leu Ser Tyr Gly  
165 170 175

Val Gln Cys Phe Ser Arg Tyr Pro Asp His Met Lys Arg His Asp Phe  
180 185 190

Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe  
195 200 205

Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu  
210 215 220

## PA0394 PCT - Sequence Listing - Patent in 3.2.ST25

Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys  
 225 230 235 240

Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser  
 245 250 255

His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val  
 260 265 270

Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Gly Val Gln Leu Ala  
 275 280 285

Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu  
 290 295 300

Pro Asp Asn His Tyr Leu Ser Thr Gln Ser Ala Leu Ser Lys Asp Pro  
 305 310 315 320

Asn Glu Lys Arg Asp His Met Val Leu Leu Gly Phe Val Thr Ala Ala  
 325 330 335

Gly Ile Thr His Gly Met Asp Glu Leu Tyr Lys  
 340 345

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<210> 13  
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## PA0394 PCT - Sequence Listing - Patentin 3.2.ST25

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